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Raw Sequence Listing Error Summary

01PE

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/072,809

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 ✓ Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



**Does Not Comply
Corrected Diskette Needed**

OIKE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/072,809

DATE: 05/24/2002

TIME: 12:06:02

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05242002\J072809.raw

3 <110> APPLICANT: Anderson, Marilyn, A., Lay, Fung T., Heath, Robyn, L.
 5 <120> TITLE OF INVENTION: Plant-derived molecules and genetic sequences encoding same
 and uses
 6 therefor
 8 <130> FILE REFERENCE: 18-01
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/072,809
 C--> 11 <141> CURRENT FILING DATE: 2002-05-01
 13 <150> PRIOR APPLICATION NUMBER: USSN 60/267,271
 14 <151> PRIOR FILING DATE: 2001-02-08
 16 <160> NUMBER OF SEQ ID NOS: 61
 18 <170> SOFTWARE: PatentIn version 3.0
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 28
 22 <212> TYPE: DNA
 23 <213> ORGANISM: primer
 25 <400> SEQUENCE: 1
 26 ggaattccat atggctcgct ccttgctg 28
 28 <210> SEQ ID NO: 2
 29 <211> LENGTH: 29
 30 <212> TYPE: DNA
 31 <213> ORGANISM: primer
 33 <400> SEQUENCE: 2
 34 gcggatcctc agttatccat tatctcttc 29
 36 <210> SEQ ID NO: 3
 37 <211> LENGTH: 24
 38 <212> TYPE: DNA
 39 <213> ORGANISM: primer
 41 <400> SEQUENCE: 3
 42 ccggatccag agaatgcaaa acag 24
 44 <210> SEQ ID NO: 4
 45 <211> LENGTH: 26
 46 <212> TYPE: DNA
 47 <213> ORGANISM: primer
 49 <400> SEQUENCE: 4
 50 gggagctctt agttatccat tatctc 26
 52 <210> SEQ ID NO: 5
 53 <211> LENGTH: 31
 54 <212> TYPE: DNA
 55 <213> ORGANISM: primer
 57 <400> SEQUENCE: 5
 58 ggaattctaa acaatggctc gctccttgctg c 31
 60 <210> SEQ ID NO: 6
 61 <211> LENGTH: 29
 62 <212> TYPE: DNA

invalid response, see error summary sheet item 10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/072,809

DATE: 05/24/2002

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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05242002\J072809.raw

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63 <213> ORGANISM: primer
65 <400> SEQUENCE: 6
66 gctctagatc agttatccat tatctcttc 29
68 <210> SEQ ID NO: 7
69 <211> LENGTH: 141
70 <212> TYPE: DNA
71 <213> ORGANISM: Nicotiana alata
73 <220> FEATURE:
74 <221> NAME/KEY: CDS
75 <222> LOCATION: (1)..(141)
77 <400> SEQUENCE: 7
78 aga gaa tgc aaa aca gaa agc aac aca ttt cct gga ata tgc att acc 48
79 Arg Glu Cys Lys Thr Glu Ser Asn Thr Phe Pro Gly Ile Cys Ile Thr
80 1 5 10 15
82 aaa cca cca tgc aga aaa gct tgt atc agt gag aaa ttt act gat ggt 96
83 Lys Pro Pro Cys Arg Lys Ala Cys Ile Ser Glu Lys Phe Thr Asp Gly
84 20 25 30
86 cat tgt agc aaa atc ctc aga agg tgc cta tgt act aag cca tgt 141
87 His Cys Ser Lys Ile Leu Arg Arg Cys Leu Cys Thr Lys Pro Cys
88 35 40 45
90 <210> SEQ ID NO: 8
91 <211> LENGTH: 47
92 <212> TYPE: PRT
93 <213> ORGANISM: Nicotiana alata
95 <400> SEQUENCE: 8
96 Arg Glu Cys Lys Thr Glu Ser Asn Thr Phe Pro Gly Ile Cys Ile Thr
97 1 5 10 15
99 Lys Pro Pro Cys Arg Lys Ala Cys Ile Ser Glu Lys Phe Thr Asp Gly
100 20 25 30
102 His Cys Ser Lys Ile Leu Arg Arg Cys Leu Cys Thr Lys Pro Cys
103 35 40 45
105 <210> SEQ ID NO: 9
106 <211> LENGTH: 75
107 <212> TYPE: DNA
108 <213> ORGANISM: Nicotiana alata
110 <220> FEATURE:
111 <221> NAME/KEY: CDS
112 <222> LOCATION: (1)..(75)
114 <400> SEQUENCE: 9
115 atg gct cgc tcc ttg tgc ttc atg gca ttt gct atc ttg gca agg atg 48
116 Met Ala Arg Ser Leu Cys Phe Met Ala Phe Ala Ile Leu Ala Arg Met
117 1 5 10 15
119 ctc ttt gtt gcc tat gag gtg caa gct 75
120 Leu Phe Val Ala Tyr Glu Val Gln Ala
121 20 25
123 <210> SEQ ID NO: 10
124 <211> LENGTH: 25
125 <212> TYPE: PRT
126 <213> ORGANISM: Nicotiana alata

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/072,809

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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05242002\J072809.raw

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128 <400> SEQUENCE: 10
129 Met Ala Arg Ser Leu Cys Phe Met Ala Phe Ala Ile Leu Ala Arg Met
130 1 5 10 15
132 Leu Phe Val Ala Tyr Glu Val Gln Ala
133 20 25
135 <210> SEQ ID NO: 11
136 <211> LENGTH: 99
137 <212> TYPE: DNA
138 <213> ORGANISM: Nicotiana alata
140 <220> FEATURE:
141 <221> NAME/KEY: CDS
142 <222> LOCATION: (1)..(99)
144 <400> SEQUENCE: 11
145 gtg ttt gat gag aag atg act aaa aca gga gct gaa att ttg gct gag 48
146 Val Phe Asp Glu Lys Met Thr Lys Thr Gly Ala Glu Ile Leu Ala Glu
147 1 5 10 15
149 gaa gca aaa act ttg gct gca gct ttg ctt gaa gaa gag ata atg gat 96
150 Glu Ala Lys Thr Leu Ala Ala Ala Leu Leu Glu Glu Glu Ile Met Asp
151 20 25 30
153 aac 99
154 Asn
156 <210> SEQ ID NO: 12
157 <211> LENGTH: 33
158 <212> TYPE: PRT
159 <213> ORGANISM: Nicotiana alata
161 <400> SEQUENCE: 12
162 Val Phe Asp Glu Lys Met Thr Lys Thr Gly Ala Glu Ile Leu Ala Glu
163 1 5 10 15
165 Glu Ala Lys Thr Leu Ala Ala Ala Leu Leu Glu Glu Glu Ile Met Asp
166 20 25 30
168 Asn
170 <210> SEQ ID NO: 13
171 <211> LENGTH: 216
172 <212> TYPE: DNA
173 <213> ORGANISM: Nicotiana alata
175 <220> FEATURE:
176 <221> NAME/KEY: CDS
177 <222> LOCATION: (1)..(216)
179 <400> SEQUENCE: 13
180 atg gct cgc tcc ttg tgc ttc atg gca ttt gct atc ttg gca agg atg 48
181 Met Ala Arg Ser Leu Cys Phe Met Ala Phe Ala Ile Leu Ala Arg Met
182 1 5 10 15
184 ctc ttt gtt gcc tat gag gtg caa gct aga gaa tgc aaa aca gaa agc 96
185 Leu Phe Val Ala Tyr Glu Val Gln Ala Arg Glu Cys Lys Thr Glu Ser
186 20 25 30
188 aac aca ttt cct gga ata tgc att acc aaa cca cca tgc aga aaa gct 144
189 Asn Thr Phe Pro Gly Ile Cys Ile Thr Lys Pro Pro Cys Arg Lys Ala
190 35 40 45
192 tgt atc agt gag aaa ttt act gat ggt cat tgt agc aaa atc ctc aga 192

```

RAW SEQUENCE LISTING

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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05242002\J072809.raw

```

193 Cys Ile Ser Glu Lys Phe Thr Asp Gly His Cys Ser Lys Ile Leu Arg
194      50                      55                      60
196 agg tgc cta tgt act aag cca tgt
197 Arg Cys Leu Cys Thr Lys Pro Cys
198 65                      70
200 <210> SEQ ID NO: 14
201 <211> LENGTH: 72
202 <212> TYPE: PRT
203 <213> ORGANISM: Nicotiana alata
205 <400> SEQUENCE: 14
206 Met Ala Arg Ser Leu Cys Phe Met Ala Phe Ala Ile Leu Ala Arg Met
207 1                      5                      10                      15
209 Leu Phe Val Ala Tyr Glu Val Gln Ala Arg Glu Cys Lys Thr Glu Ser
210      20                      25                      30
212 Asn Thr Phe Pro Gly Ile Cys Ile Thr Lys Pro Pro Cys Arg Lys Ala
213      35                      40                      45
215 Cys Ile Ser Glu Lys Phe Thr Asp Gly His Cys Ser Lys Ile Leu Arg
216      50                      55                      60
218 Arg Cys Leu Cys Thr Lys Pro Cys
219 65                      70
221 <210> SEQ ID NO: 15
222 <211> LENGTH: 240
223 <212> TYPE: DNA
224 <213> ORGANISM: Nicotiana alata
226 <220> FEATURE:
227 <221> NAME/KEY: CDS
228 <222> LOCATION: (1)..(240)
230 <400> SEQUENCE: 15
231 aga gaa tgc aaa aca gaa agc aac aca ttt cct gga ata tgc att acc
232 Arg Glu Cys Lys Thr Glu Ser Asn Thr Phe Pro Gly Ile Cys Ile Thr
233 1                      5                      10                      15
235 aaa cca cca tgc aga aaa gct tgt atc agt gag aaa ttt act gat ggt
236 Lys Pro Pro Cys Arg Lys Ala Cys Ile Ser Glu Lys Phe Thr Asp Gly
237      20                      25                      30
239 cat tgt agc aaa atc ctc aga agg tgc cta tgt act aag cca tgt gtg
240 His Cys Ser Lys Ile Leu Arg Arg Cys Leu Cys Thr Lys Pro Cys Val
241      35                      40                      45
243 ttt gat gag aag atg act aaa aca gga gct gaa att ttg gct gag gaa
244 Phe Asp Glu Lys Met Thr Lys Thr Gly Ala Glu Ile Leu Ala Glu Glu
245      50                      55                      60
247 gca aaa act ttg gct gca gct ttg ctt gaa gaa gag ata atg gat aac
248 Ala Lys Thr Leu Ala Ala Leu Leu Glu Glu Glu Ile Met Asp Asn
249 65                      70                      75                      80
251 <210> SEQ ID NO: 16
252 <211> LENGTH: 80
253 <212> TYPE: PRT
254 <213> ORGANISM: Nicotiana alata
256 <400> SEQUENCE: 16
257 Arg Glu Cys Lys Thr Glu Ser Asn Thr Phe Pro Gly Ile Cys Ile Thr

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/072,809

DATE: 05/24/2002

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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05242002\J072809.raw

```

258 1          5          10          15
260 Lys Pro Pro Cys Arg Lys Ala Cys Ile Ser Glu Lys Phe Thr Asp Gly
261          20          25          30
263 His Cys Ser Lys Ile Leu Arg Arg Cys Leu Cys Thr Lys Pro Cys Val
264          35          40          45
266 Phe Asp Glu Lys Met Thr Lys Thr Gly Ala Glu Ile Leu Ala Glu Glu
267          50          55          60
269 Ala Lys Thr Leu Ala Ala Ala Leu Leu Glu Glu Glu Ile Met Asp Asn
270 65          70          75          80
272 <210> SEQ ID NO: 17
273 <211> LENGTH: 541
274 <212> TYPE: DNA
275 <213> ORGANISM: Nicotiana alata
277 <220> FEATURE:
278 <221> NAME/KEY: CDS
279 <222> LOCATION: (1)..(318)
281 <400> SEQUENCE: 17
282 atg gct cgc tcc ttg tgc ttc atg gca ttt gct atc ttg gca agg atg      48
283 Met Ala Arg Ser Leu Cys Phe Met Ala Phe Ala Ile Leu Ala Arg Met
284 1          5          10          15
286 ctc ttt gtt gcc tat gag gtg caa gct aga gaa tgc aaa aca gaa agc      96
287 Leu Phe Val Ala Tyr Glu Val Gln Ala Arg Glu Cys Lys Thr Glu Ser
288          20          25          30
290 aac aca ttt cct gga ata tgc att acc aaa cca cca tgc aga aaa gct      144
291 Asn Thr Phe Pro Gly Ile Cys Ile Thr Lys Pro Pro Cys Arg Lys Ala
292          35          40          45
294 tgt atc agt gag aaa ttt act gat ggt cat tgt agc aaa atc ctc aga      192
295 Cys Ile Ser Glu Lys Phe Thr Asp Gly His Cys Ser Lys Ile Leu Arg
296          50          55          60
298 agg tgc cta tgt act aag cca tgt gtg ttt gat gag aag atg act aaa      240
299 Arg Cys Leu Cys Thr Lys Pro Cys Val Phe Asp Glu Lys Met Thr Lys
300 65          70          75          80
302 aca gga gct gaa att ttg gct gag gaa gca aaa act ttg gct gca gct      288
303 Thr Gly Ala Glu Ile Leu Ala Glu Glu Ala Lys Thr Leu Ala Ala Ala
304          85          90          95
306 ttg ctt gaa gaa gag ata atg gat aac taa ttagagatta gaagaaatta      338
307 Leu Leu Glu Glu Glu Ile Met Asp Asn
308          100          105
310 aggatgcagt atcacacata ataaagtttc tacctttctt aaaagtgtag ctaatgttgt      398
312 gttttaattg gcttttagta gccttttatt acactttaaa taagtgtggc acttcaatcc      458
314 tttgtgcaat cttgcactaa gtttatttgt gtacttttaa tgaaaatgac cttctatggt      518
316 ctttggttaa aaaaaaaaaa aaa      541
318 <210> SEQ ID NO: 18
319 <211> LENGTH: 105
320 <212> TYPE: PRT
321 <213> ORGANISM: Nicotiana alata
323 <400> SEQUENCE: 18
324 Met Ala Arg Ser Leu Cys Phe Met Ala Phe Ala Ile Leu Ala Arg Met
325 1          5          10          15

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/072,809

DATE: 05/24/2002
TIME: 12:06:03

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05242002\J072809.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:58; Xaa Pos. 1,2,4,5,6,8,9,11,13,15,16,17,18,19,21,22,23,25,26,28,30
Seq#:58; Xaa Pos. 31,33,35,36,37,38,39,40,42,45,46
Seq#:59; Xaa Pos. 2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23
Seq#:59; Xaa Pos. 24,25,26,27,28,29,30,32
Seq#:60; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22
Seq#:60; Xaa Pos. 23,24,25,26,27,28,29,30,31,32,33
Seq#:61; Xaa Pos. 2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23
Seq#:61; Xaa Pos. 24,25,26,27,28,29,30,32,33,34,36,37,38,40,41,43,45,47,48
Seq#:61; Xaa Pos. 49,50,51,53,54,55,57,58,60,62,63,65,67,68,69,70,71,72,74
Seq#:61; Xaa Pos. 77,78,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96
Seq#:61; Xaa Pos. 97,98,99,100,101,102,103,104,105,106,107,108,109,110,111
Seq#:61; Xaa Pos. 112

VERIFICATION SUMMARY

DATE: 05/24/2002

PATENT APPLICATION: US/10/072,809

TIME: 12:06:03

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05242002\J072809.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1348 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:0
L:1351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:16
L:1354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:32
L:1508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
L:1511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:16
L:1685 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:0
L:1688 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:16
L:1691 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:32
L:2169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0
L:2172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:16
L:2175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:32
L:2178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:48
L:2181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:64
L:2184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:80
L:2187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:96